WHAT IS CLAIMED IS:

- 1. A polypeptide capable of modulating the autoimmune response of an individual to acetylcholine receptor, said polypeptide being selected from the group consisting of:
- (i) a polypeptide consisting of the amino acid sequence of SEQ ID NO:6;
- (ii) a polypeptide consisting of the amino acid sequence of SEQ ID NO:8;
- (iii) a polypeptide corresponding to amino acid residues 1-121 of SEQ ID NO:2;
- (iv) a polypeptide corresponding to amino acid residues 1-146 of SEQ ID NO:6;
- (v) a polypeptide corresponding to amino acid residues 122-210 of SEQ ID NO:2;
- (vi) a polypeptide as in (i) to (v) or the polypeptide $H\alpha 1-210$ of SEQ ID NO:2 in which one or more amino acid residues have been added, deleted or substituted by other amino acid residues in a manner that the resulting polypeptide is capable of suppressing experimental myasthenia gravis in animal models;
- (vii) a fragment of a polypeptide as in (i) to (vi),
 which fragment is capable of suppressing experimental
 myasthenia gravis in animal models;

The paper and the first the paper than the paper th

(viii) a polypeptide comprising two or more
fragments as in (vii) fused together with or without a spacer;

- (ix) a polypeptide, or a fragment as defined in (i)- (viii), or the polypeptide H α 1-210 of SEQ ID NO:2, fused to an additional polypeptide at its N- and/or C-termini; and
- (x) soluble forms, denatured forms, chemical derivatives and salts of a polypeptide or a fragment as defined in (i)-(ix).
- 2. A polypeptide according to claim 1, wherein said polypeptide consists of the amino acid sequence of SEQ ID NO:6.
- 3. A polypeptide according to claim 1, wherein said polypeptide consists of the amino acid sequence of SEQ ID ${\tt NO:8}$.
- 4. A polypeptide according to claim 1, corresponding to amino acid residues 1-121 of SEQ ID NO:2.
- 5. A polypeptide according to claim 1, corresponding to amino acid residues 1-146 of SEQ ID NO:6.
- 6. A polypeptide according to claim 1, corresponding to amino acid residues 122-210 of SEQ ID NO:2.
- 7. A polypeptide according to claim 1, wherein an additional polypeptide, which is glutathione S-transferase (GST), is fused to said polypeptide or fragment thereof at the N-terminus of said polypeptide or fragment thereof.

- 8. A DNA molecule coding for the polypeptide according to claim 1.
- 9. A DNA molecule according to claim 8, being selected from the group consisting of:
- (i) a DNA molecule comprising the nucleotide sequence of SEQ ID NO:5;
- (ii) a DNA molecule comprising the nucleotide
 sequence of SEQ ID NO:7;
- (iii) a DNA molecule comprising the nucleotide corresponding to nucleotides 1 to 363 of SEQ ID NO:1;
- (iv) a DNA molecule comprising the nucleotide sequence corresponding nucleotides 1 to 438 of SEQ ID NO:5;
- (v) a DNA molecule comprising the nucleotide sequence of nucleotides 364 to 630 of SEQ ID NO:1;
- (vi) DNA molecules which are degenerate, as a result of the genetic code, to the DNA sequences of (i) to (v) and which code for a polypeptide coded for by any one of the DNA sequences of (i) to (v);
- (vii) a DNA molecule having a coding nucleotide sequence which is at least 70% homologous to any one of the DNA sequences of (i) to (vi) or to the DNA sequence, SEQ ID NO:1, coding for $H\alpha 1-210$;
- (viii) a DNA molecule as in (i) to (v) or the DNA molecule coding for the amino acid sequence SEQ ID NO:2 of

I H Hert and the E H West the time that the first there had the the

 $H\alpha 1-210$, in which one or more codons has been added, replaced or deleted in a manner that the polypeptide coded for by said sequence is capable of suppressing experimental myasthenia gravis in animal models;

- (ix) a fragment of a DNA molecule as in (i)-(viii)
 which codes for a polypeptide capable of suppressing
 experimental myasthenia gravis in animal models;
- (x) a DNA molecule comprising two or more fragments of (ix) fused together with or without a spacer, and which codes for a polypeptide capable of suppressing experimental myasthenia gravis in animal models; and
- (xi) a DNA molecule comprising a nucleic acid sequence as defined in (i)-(x) or the DNA sequence, SEQ ID NO:1, coding for H α 1-210, fused to additional coding DNA sequences at its 3' and/or 5' end.
- 10. A DNA molecule according to claim 9, which comprises the nucleotide sequence of SEQ ID NO:5.
- 11. A DNA molecule according to claim 9, which comprises the nucleotide sequence of SEQ ID NO:7.
- 12. A DNA molecule according to claim 9, which comprises the nucleotide sequence corresponding to nucleotides 1 to 363 of SEQ ID NO:1.

- 13. A DNA molecule according to claim 9, which comprises the nucleotide sequence of nucleotides 1 to 438 of SEO ID NO:5.
- 14. A DNA molecule according to claim 9, which comprises the nucleotide sequence of nucleotides 364 to 630 of SEQ ID NO:1.
- 15. A DNA molecule according to claim 9, wherein said additional coding sequence in (xi) codes for glutathione S-transferase (GST) and is fused at the 5' end of said nucleic acid sequence.
- 16. A replicable expression vehicle comprising a DNA molecule according to claim 8.
- 17. A prokaryotic or eukaryotic host cell transformed with the replicable expression vehicle of claim 16.
- 18. A process for preparing a polypeptide capable of modulating the autoimmune response of an individual to acetylcholine receptor, comprising:
- (i) culturing a host cell of claim 17 under conditions promoting expression; and
 - (ii) isolating the expressed polypeptide.
- 19. A process according to claim 18, wherein the expressed polypeptide is a fused polypeptide.

- 20. A pharmaceutical composition, comprising a pharmaceutically acceptable carrier and the polypeptide of claim 1 or a polypeptide having the amino acid sequence of SEQ ID NO:2.
- 21. A method for alleviating and/or treating myasthenia gravis, comprising administering to an individual in need thereof an effective amount of a polypeptide according to claim 1 or of a polypeptide having the amino acid sequence of SEQ ID NO:2.
- 22. A method for diagnosing myasthenia gravis, comprising:
- (i) incubating one or more polypeptides selected from the group consisting of (i) to (x) of claim 1, and a polypepetide having the amino acid sequence of SEQ ID NO:2;
- (ii) determining the amount of the anti-AChR antibodies in the serum bound to said one or more polypeptides,

whereby detection of anti-AChR titers indicates the presence of myasthenia gravis.